

ERRATUM

Open Access



Erratum to: Functional signatures of oral dysbiosis during periodontitis progression revealed by microbial metatranscriptome analysis

Susan Yost^{1†}, Ana E Duran-Pinedo^{1†}, Ricardo Teles³, Keerthana Krishnan¹ and Jorge Frias-Lopez^{1,2*}

Erratum

It has come to our attention that there is an error in the methods section of this article [1]. The sentence: “Quantified libraries were pooled and sequenced using the MiSeq v2, 2 × 150 cycle cartridge (Illumina)” is incorrect and should read: “Quantified libraries were pooled and sequenced using the MiSeq v3, 2 × 75 cycle cartridge (Illumina)”. The publisher apologises for any inconvenience caused.

Author details

¹Forsyth Institute, 245 First Street, Cambridge 02142, Massachusetts, USA.

²Harvard School of Dental Medicine, 188 Longwood Ave, Boston 02115, MA, USA. ³University of North Carolina Chapel Hill, School of Dentistry, Chapel Hill 27599-7450, NC, USA.

Published online: 27 October 2015

Reference

1. Yost S, Duran-Pinedo AE, Teles R, Krishnan K, Frias-Lopez J. Functional signatures of oral dysbiosis during periodontitis progression revealed by microbial metatranscriptome analysis. *Genome Med.* 2015;7:27.

* Correspondence: jfrias@forsyth.org

†Equal contributors

¹Forsyth Institute, 245 First Street, Cambridge 02142, Massachusetts, USA

²Harvard School of Dental Medicine, 188 Longwood Ave, Boston 02115, MA, USA

Full list of author information is available at the end of the article